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STOP ATG ATG

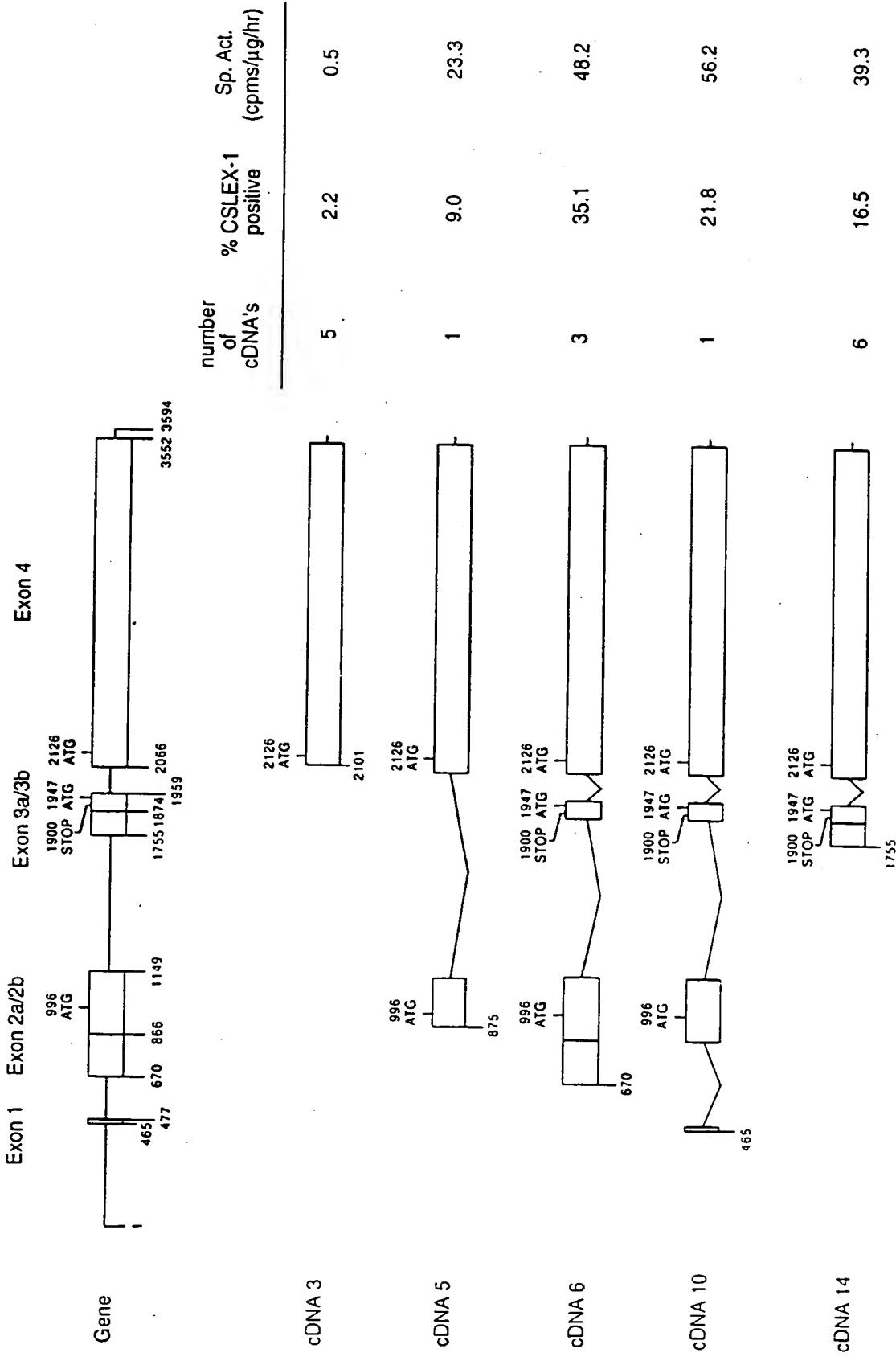


FIGURE 1a

112 — 74 — 45 — 30 —

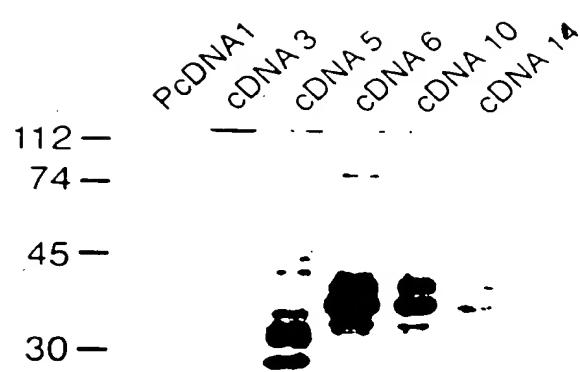


FIGURE 1b

2

acaaacaggaaggaca~~g~~caggctctggcagccagaagcctgtggcccaagctggcaggatggcccttcgcaggtc 80  
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 cctgggtcaccagcttgggagaggtgaagaaagatgccggggcggaaacaaaggggcagatatactatggttatctt 240  
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 ggccaaacactgtcctccattcaaaactcCCAGTCTGGAGCTCTGGACATGGACAAGCCAGGCCTGCTATTCTCCATAC 720  
 AGGGCTCCATAGTGTCTGGCTCAGCAGAGTGGGGATCTGGTGGGATGGAGGAAGCTTAGCTAAAGCTTGTATAGGC 800  
 TGAAGCTCTGAGTGACCTGCTGGCCACCCCTACCCCTGGTCTGGCTGGTCATTGCATCCCCAGATMGAAGGCTTGGT 880  
 GAGATGGAGAGGAACCTTGGCTACAAGCTATAGCTTGCACCAGAGCCTGCTGGAGGGGAATCAAACAAGCCTGGACC 960  
 TGAGGCTGGACTAGCTTCCTGTTCTGGAGTGGATGCCAACCCCTGCCACCAGCCTGCTGGAGGGACA 1040  
 M P T P C P P A C L S T P G T  
 CACAGACTCCTCCCTTCCAGACTGGAAAGCCCCCTCCTGGAGAGCAGGAAGGAAGCAACCTGCAACTCTCCAGGCC 1120  
 H R L L P F P D W K A P S W E S R K E A T C N S S S P  
 TGGACCTTGGCCTGAACCTACAGTTCAAGgttgtatgctcacaggcttggcaggaaagataagaatccccagggcac 1200  
 G P W A E P T V Q  
 cctcccccccccagtcactgcaggttagctcctgggtctgccttcaggcaagtgcgtacgcgtccatcagactg 1280  
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 TGCCAAAGATTGAAGGGTAGGGGGCAGAAGTGGGAAGGTCCCTGGCTTCACCTGGTAGATGGTGGAGGCC 1840  
 CAGAGGTTGAGCTGAGCAGCAGCTGTGATTTCAGGGTGCCCTGTGGAGAGGCTGCTGTGATTGAAAATCTTCTTCC 1920  
 TTGGTGACAATTCCAGAAGGCTCAGATGAATTGTATTGgtgagtgcctggcccttaagcagtcccagctgggatgatg 2000  
 M N C I

FIGURE 2

gggatttatgggtgtccctgagcctagggtagacagggccttccttttttttattctgcttcagGGTACCACCCCACC 2080  
 G Y H P T

AGGAGGCTCGGGCCTGGGGCGGCCTAGCTGGAGGAGCAACATTCACTGGTAATTGGTTTCTGGCTGTGGGATCAGC 2160  
 R R L R A W G G L A G G A T F M V I W F F W L W G S A

TCCTGGAAGTGCCTCTGTGCTCAGTCCACACTCACCATCCTTATCTGGCACTGGCTTACCAACCGGCCAGAGC 2240  
 P G S A P V P Q S T L T I L I W H W P F T N R P P E

TACCTGGTACACCTGCACTCGCTATGGCATGCCAGCTGCCGCTGAGTGCTAACCGGAGCCTGCTAGCCAGTGCTGAT 2320  
 L P G D T C T R Y G M A S C R L S A N R S L L A S A D

GCTGTGGTCTTCCACCAACCGTGAGCTGCAAACCCGGCAATCTCTCCACCCCTGGACCAGAGGCCACACGGACAGCCTG 2400  
 A V V F H H R E L Q T R Q S L L P L D Q R P H G Q P W

GGTCTGGGCCTCCATGGAATGCCAGTAATACCCATGGTCTCCATGCCCTGGGGCATCTCACTGGGTGCTGAGCT 2480  
 V W A S M E S P S N T H G L H R F R G I F N W V L S

ATCGGCGTGATTCAAGATATCTTGACCGCTACGGTCGCTTGGAGCCTCTCTGGGCCACATCCCCACTACCGGCCAAA 2560  
 Y R R D S D I F V P Y G R L E P L S G P T S P L P A K

AGCAGGATGGCTGCCCTGGGTGATCAGCAATTCCAGGAGCGGCAGCAGCGTCAAAGCTGTACCGGAGCTGGCCCTCA 2640  
 S R M A A W V I S N F Q E R Q Q R A K L Y R Q L A P H

TCTGCAGGTGGATGTGTTGGCGCCAGGGACGGCCCTATGCGTAATTGTCCTGCCCACCTTGGCCCGTAC 2720  
 L Q V D V F G R A S G R P L C A N C L L P T L A R Y

GCTTCTACCTGGCTTGAGAACTCACAGCATGGGACTACATCACTGAGAAGTTCTGGCGCAATGCCCTGGCGCTGGT 2800  
 R F Y L A F E N S Q H R D Y I T E K F W R N A L A A G

GCTGTACCCGTGGCGCTGGACCTCCCTGGGCCACCTACGAGGCTTGTGCCACAGATGCCCTGGCGACG 2880  
 A V P V A L G P P R A T Y E A F V P P D A F V H V D D

CTTCAGCTCTGCCGTGAACCTGGCTCTCGTCAGCATGAATGAGAGTCGTTATGTCCTGGCTTGTACACGTGGACGA 2960  
 F S S A R E L A V F L V S M N E S R Y R G F F A W R

ACCGGCTCCGTGCGGCTCTGGGTGACTGGAGGGAGCGCTCTGCACCATCTGTGCCCGTACCCCTACTGGCCCGC 3040  
 D R L R V R L L G D W R E R F C T I C A R Y P Y L P R

AGCCAGGTCTATGAAGACCTTGAAAGCTGGTCCAGGCTGAACTCCCTGCTGGAGAGGGCTGGATGGTGGAGACT 3120  
 S Q V Y E D L E S W F Q A \*

GATGTTGAAACCAAAGAGCTGGCATCCAGGCTTGTGACCATGGCACTACCCAAAGGCTTCTGTTAGTCAGTGAGCA 3200

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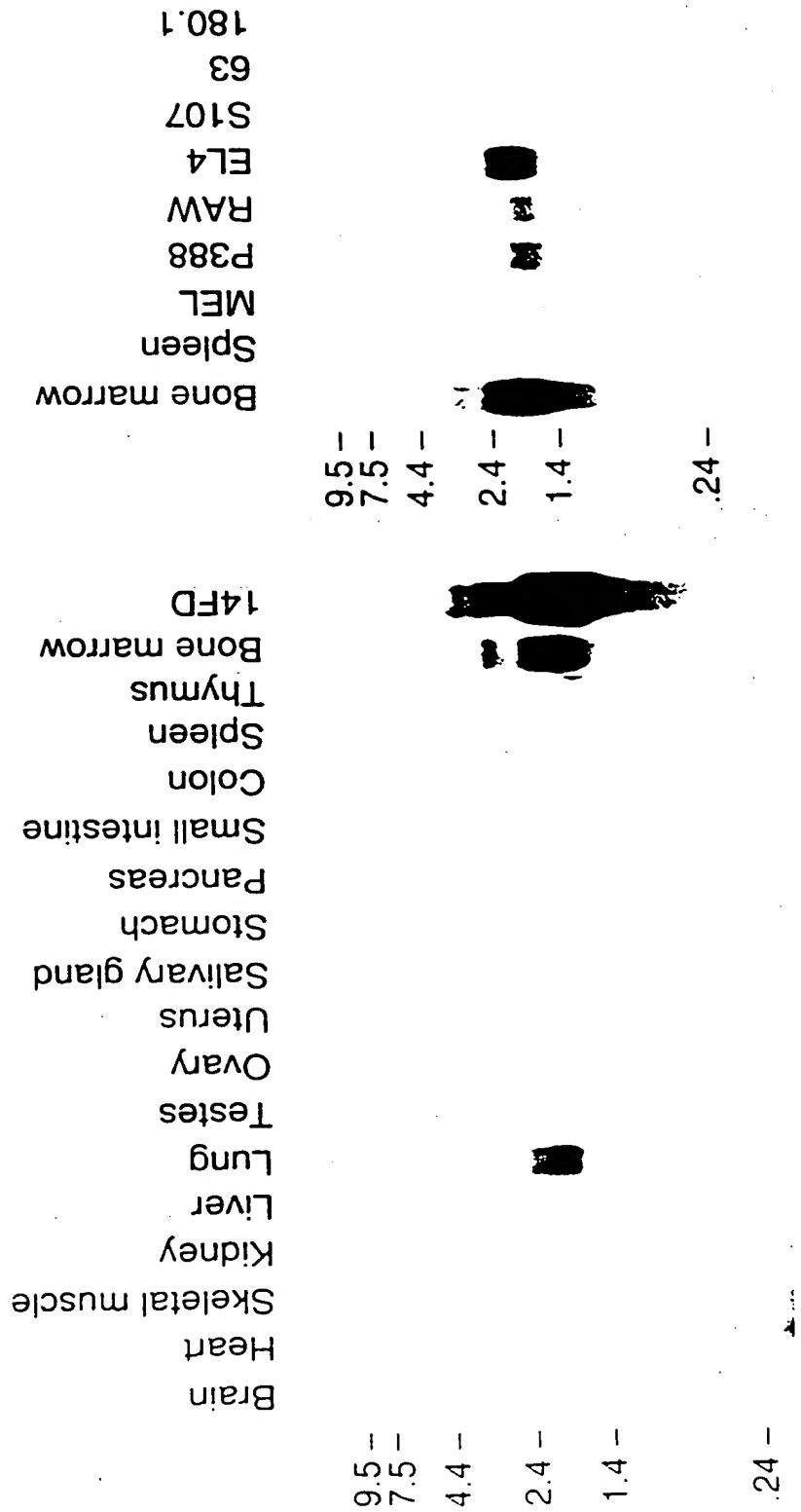
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CAGACTGTACACTGGCGCAGGGGTTGTGAAGGACAGTGCAGATGATTCTGGCTTGTACCCACAGTTCCCCCAGG 3520

GAAAGAGGCACTACTAATAAAACACTGACAGaaatccctggtaagtctgttaggcagcagagctcgaaattc 3594

FIGURE 2 - cont.



(a)

(b)

FIGURES 3a and 3b

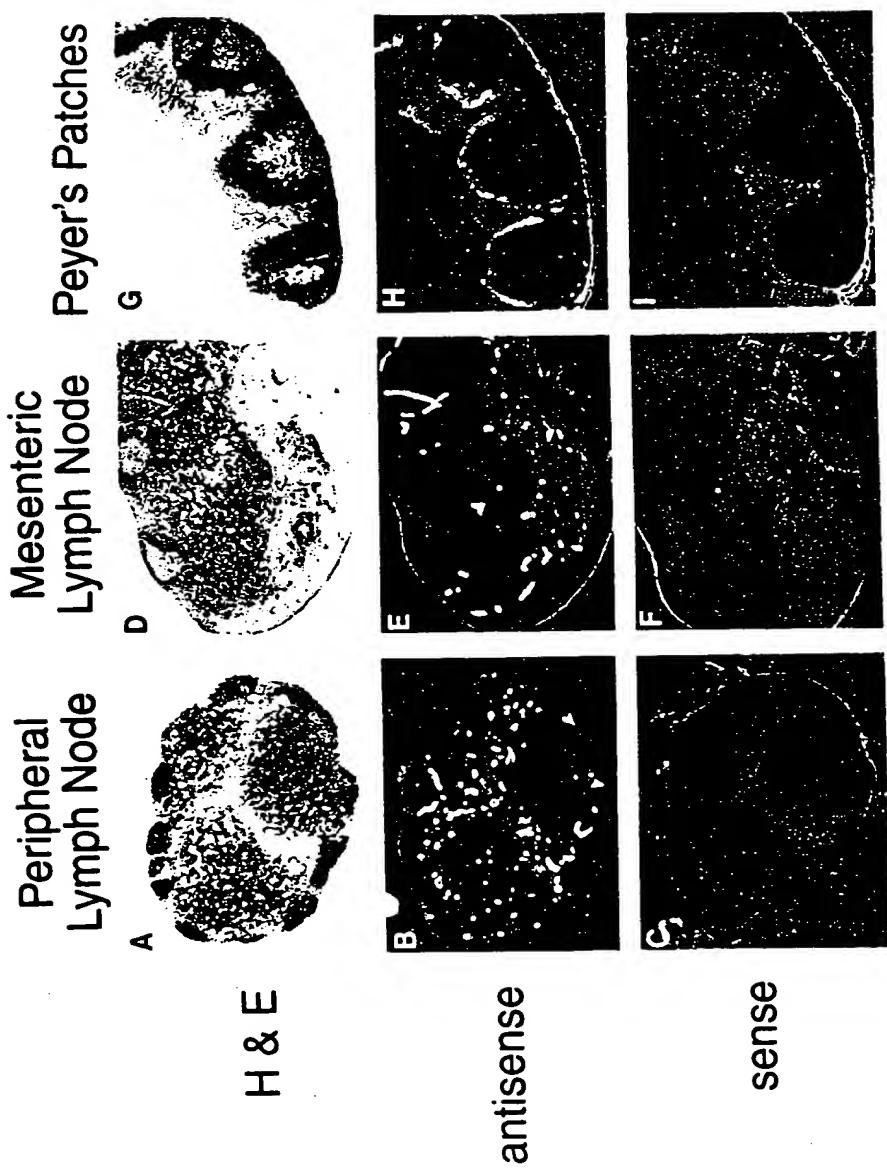
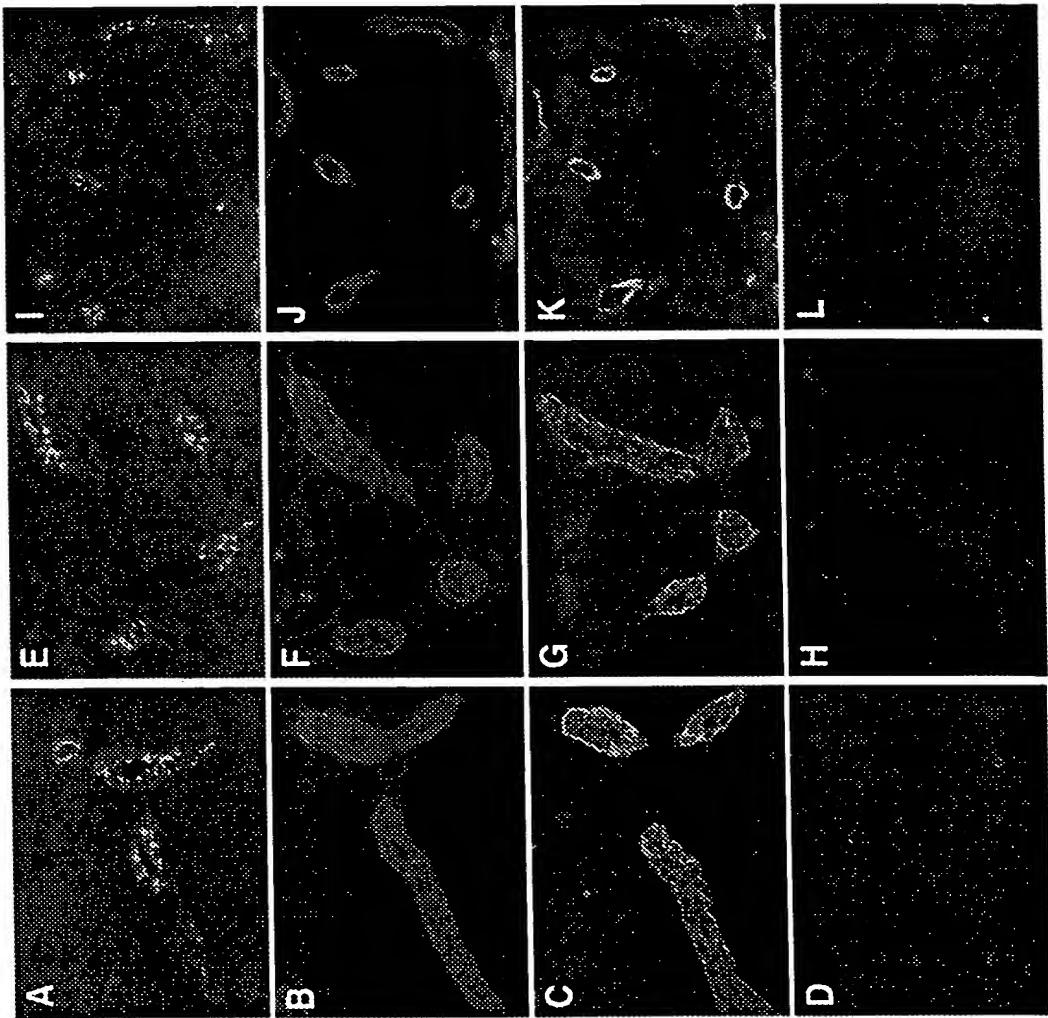


FIGURE 4

Peripheral Mesenteric Lymph Node Patches Peyer's Patches



Fuc-TVII

MECA-79

L-sel/IgM Chimera

L-sel/IgM Chimera + EDTA

FIGURE 5a

FIGURE 5b

